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SERIAL NUMBER	FILING DATE	FIRST NAMED APPLICANT	ATTORNEY DOCKET NO.
09/328,738	08/13/99	Heller, Michael J.	241/172

EXAMINER	
Thomas Prasthofer	
ART UNIT	PAPER NUMBER
1627	16

Please find below a communication from the EXAMINER in charge of this application

~~NON-BONA FIDE ATTEMPT to Comply with Sequence Rules~~

This application contains sequence disclosures that are encompassed by the definitions for nucleotide and/or amino acid sequences set forth in 37 C.F.R. § 1.821(a)(1) and (a)(2). However, this application fails to comply with the requirements of 37 C.F.R. §§ 1.821-1.825 for the reason(s) set forth on the attached Notice To Comply With Requirements For Patent Applications Containing Nucleotide Sequence And/Or Amino Acid Sequence Disclosures.

APPLICANT IS GIVEN 30 days FROM THE DATE OF THIS LETTER WITHIN WHICH TO COMPLY WITH THE SEQUENCE RULES, 37 C.R.F. §§ 1.821-1.825. Failure to comply with these requirements will result in ABANDONMENT of the application under 37 C.F.R. § 1.821(g). Extensions of time may be obtained by filing a petition accompanied by the extension fee under the provisions of 37 C.F.R. § 1.136. In no case may an applicant extend the period for response beyond the six month statutory period. Direct the response to the undersigned. Applicant is requested to return a copy of the attached Notice to Comply with the response.

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Examiner, whose telephone number is (703) 308-4548. If the examiner cannot be reached, inquiries can be directed to Supervisory Patent Examiner, whose telephone number is (703) 308-2439. The fax number for the organization where this application or proceeding is assigned is (703) 308-2742.

Any inquiry of a general nature or relating to the status of this application should be directed to the Group receptionist whose telephone number is (703) 308-1235.

for SAE
for
PADMA SHRI PONNALURI
PRIMARY EXAMINER



filename: 09374338.prj

CProject Generation cancelled by user.~

CProjectData 3 MICROELECTRONIC MOLECULAR DESCRIPTOR ARRAY DEVICES, METHODS, PROCEDURES, AND FORMATS FOR COMBINATORIAL SELECTION OF INTERMOLECULAR LIGAND BINDING STRUCTURES AND FOR DRUG SCREENING" Patrick Bagleyan; Nunojoh
241/17209/374, 3381999-08-13 C:\My Sequences\Sequences ~

CDNASequence 01 SYNTHETIC CONSTRUCTIONseqggg
? agctt cymkswbdhvn DNA

OtherFeature

modified_base Entire sequence is Pyranosyl RNA ? ! ?

modified_base Base 1 is tryptamine 1 1 ?

modified_base Base 7 is modified with Texas

Res 7 ? 02 SYNTHETIC CONSTRUCTIONseqccccc
? agctt cymkswbdhvn DNA ?

modified_base Entire sequence is Pyranosyl RNA 1 1 ?

modified_base-Base 1 is modified with Biotin 1 1 ?

modified

sample of submitted file

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